SEQUENCE LISTING

<110> POQUET, ISABELLE
      LLULL, DANIEL

<120> ZINC-REGULATED PROKARYOTIC EXPRESSION CASSETTES

<130> 40526U

<140> 10/525,449
<141> 2005-02-24

<150> PCT/FR03/02606
<151> 2003-08-29

<150> FR 02 10805
<151> 2002-08-30

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<170> PatentIn version 3.5

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57

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1   5   10   15

Val Ser Ser Thr Ala Val Val Phe Ala
            20   25

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<220>
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<210> 8
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35

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acaaaagctg ggtaccgggc cccccctcga ggtcgacggt atcgatagcc cgcctaatga   120

gcgggctttt ttttgatatc gaattacccg ggaattcaga tctttgatca aggatctgtc   180

agctggttca actagcggtg gtcaaactgt tagtaataaa acttattgtt ttgatgttcg   240

gcttaaggat ggaaggattt ttcaaataaa aaagtaaaaa ataatgttaa ctggttgaca   300

ttatttttac tttgctatat aattaaccag taaactaatt atggaggaca aaatactatg   360

antttagcaa atcaaatcga ccagtttctt ggggcaatta tgcagtttgc anaaaacaag   420

catgaaatat tactcggcga atgcnaaagt aatgttaagc taacaagcac gcaagaacat   480

480

atcttaatga ttctagctgc agaggtttcg acaaacgcga gaattgccga gcaactcaag
540

atttcgccag cagcggtaac taaagctctc aaaaaattac aagagcaaga actgattaaa
600

tcaagtcggg caacaaatga cgaacgcgta gtcctttgga gcctgacaga aaaagcaatt
660

ccagttgcta agaacatgc tgctcatcat gagaaaactc taagtaccta ccaagaatta
720

ggagacaaat ttactgacga agaacaaaaa gtgataagtc aattcttatc agtacttacg
780

gaggagtttc gatgaagaaa atattgatgt tatttgctat ccggcagtt ttacttcttg
840

ctggttgtca aaaaacagca gacaaaccag aagttgtgac aactttttgag ccgatgtatg
900

aatttacgaa agcgattgtt ggagataagg ttaaaattga aaatattgtt ccggcgaatc
960

aagaagttca cgaatttgaa ccgagtgcca ttacgaaaaa aatggtagaa aatgcaaaga
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1080

atatttatgc tgatggaaaa
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gatgttcggc ttaaggatgg aaggattttt caaataaaaa agtaaaaaat aatgttaact
120

ggttgacatt attttttactt tgctatataa ttaaccagta
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<220>
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caggaaacag ctatgacc
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<210> 12
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<400> 12
gttctaagga tccattaact taaggag
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<210> 13
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1   5   10   15

Val Ser Ser Thr Ala Val Val Phe Ala Tyr Ala
            20   25

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Phe Ala Glu Asn Lys His Glu Ile Leu Leu Gly Lys Cys Glu Ser Asp
            20   25   30

Val Lys Leu Thr Ser Thr Gln Glu His Ile Leu Met Leu Leu Ala Glu
        35   40   45

Gln Ile Ser Thr Asn Ala Lys Ile Ala Glu Lys Leu Lys Ile Ser Pro
    50   55   60

Ala Ala Val Thr Lys Ala Leu Lys Lys Leu Gln Glu Gln Glu Leu Ile
65   70   75   80

Lys Ser Ser Arg Ala Thr Asn Asp Glu Arg Val Val Leu Trp Ser Leu
                85   90   95

Thr Glu Lys Ala Val Pro Val Ala Lys Glu His Ala Thr His His Glu
                100   105   110

Lys Thr Leu Ser Thr Tyr Gln Glu Leu Gly Asn Lys Phe Thr Asp Glu
        115   120   125

Glu Gln Glu Val Ile Ser Lys Phe Leu Ser Ala Leu Thr Glu Glu Phe
        130   135   140

Gln
145